

SEOUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT:
 - (A) NAME: E. I. DUPONT DE NEMOURS AND COMPANY
 - (B) STREET: 1007 MARKET STREET
 - (C) CITY: WILMINGTON
 - (D) STATE: DELAWARE
 - (E) COUNTRY: U.S.A.
 - (F) POSTAL CODE (ZIP): 19898
 - (G) TELEPHONE: 302-892-8112
 - (H) TELEFAX: 302-773-0164
- (i) APPLICANT:
 - (A) NAME: GENENCOR INTERNATIONAL, INC.
 - (B) STREET: 4 CAMBRIDGE PLACE

1870 SOUTH WINTON ROAD

- (C) CITY: ROCHESTER
- (D) STATE: NEW YORK
- (E) COUNTRY: U.S.A.
- (F) POSTAL CODE (ZIP): 14618
- (G) TELEPHONE:
- (H) TELEFAX:
- (ii) TITLE OF INVENTION: BIOCONVERSION OF A FERMENTABLE
 CARBON SOURCE TO 1,3-PROPANEDIOL BY A SINGLE MICROORGANISM
- (iii) NUMBER OF SEQUENCES: 46
 - (iv) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: 3.50 INCH DISKETTE
 - (B) COMPUTER: IBM
 - (C) OPERATING SYSTEM: MICROSOFT WINDOWS 3.1
 - (D) SOFTWARE: MICROSOFT WORD 6.0
 - (v) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
 - (vi) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/440,293
 - (B) FILING DATE: MAY 12, 1995
- (vii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: LINDA AXAMETHY FLOYD
 - (B) REGISTRATION NUMBER: 33,692
 - (C) REFERENCE/DOCKET NUMBER: CR-9715-B

WO 96/35796 PCT/US96/06705

INFORMATION FOR SEQ ID NO:1: (2)

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12145 base pairs

 - (B) TYPE: nucleic acid (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GTCGACCACC	ACGGTGGTGA	CTTTAATGCC	GCTCTCATGC	AGCAGCTCGG	TGGCGGTCTC	60
AAAATTCAGG	ATGTCGCCGG	TATAGTTTTT	GATAATCAGC	AAGACGCCTT	CGCCGCCGTC	120
AATTTGCATC	GCGCATTCAA	ACATTTTGTC	CGGCGTCGGC	GAGGTGAATA	TTTCCCCCGG	180
ACAGGCGCCG	GAGAGCATGC	CCTGGCCGAT	ATAGCCGCAG	TGCATCGGTT	CATGTCCGCT	240
GCCGCCGCCG	GAGAGCAGGG	CCACCTTGCC	AGCCACCGGC	GCGTCGGTGC	GGGTCACATA	300
CAGCGGGTCC	TGATGCAGGG	TCAGCTGCGG	ATGGGCTTTA	GCCAGCCCCT	GTAATTGTTC	360
ATTCAGTACA	TCTTCAACAC	GGTTAATCAG	CTTTTTCATT	ATTCAGTGCT	CCGTTGGAGA	420
AGGTTCGATG	CCGCCTCTCT	GCTGGCGGAG	GCGGTCATCG	CGTAGGGGTA	TCGTCTGACG	480
GTGGAGCGTG	CCTGGCGATA	TGATGATTCT	GGCTGAGCGG	ACGAAAAAA	GAATGCCCCG	540
ACGATCGGGT	TTCATTACGA	AACATTGCTT	CCTGATTTTG	TTTCTTTATG	GAACGTTTTT	600
GCTGAGGATA	TGGTGAAAAT	GCGAGCTGGC	GCGCTTTTTT	TCTTCTGCCA	TAAGCGGCGG	660
TCAGGATAGC	CGGCGAAGCG	GGTGGGAAAA	AATTTTTTGC	TGATTTTCTG	CCGACTGCGG	720
GAGAAAAGGC	GGTCAAACAC	GGAGGATTGT	AAGGGCATTA	TGCGGCAAAG	GAGCGGATCG	780
GGATCGCAAT	CCTGACAGAG	ACTAGGGTTT	TTTGTTCCAA	TATGGAACGT	AAAAAATTAA	840
CCTGTGTTTC	ATATCAGAAC	AAAAAGGCGA	AAGATTTTTT	TGTTCCCTGC	CGGCCCTACA	900
GTGATCGCAC	TGCTCCGGTA	CGCTCCGTTC	AGGCCGCGCT	TCACTGGCCG	GCGCGGATAA	960
CGCCAGGGCT	CATCATGTCT	ACATGCGCAC	TTATTTGAGG	GTGAAAGGAA	TGCTAAAAGT	1020
TATTCAATCT	CCAGCCAAAT	ATCTTCAGGG	TCCTGATGCT	GCTGTTCTGT	TCGGTCAATA	1080
TGCCAAAAAC	CTGGCGGAGA	GCTTCTTCGT	CATCGCTGAC	GATTTCGTAA	TGAAGCTGGC	1140
GGGAGAGAAA	GTGGTGAATG	GCCTGCAGAG	CCACGATATT	CGCTGCCATG	CGGAACGGTT	1200
TAACGGCGAA	TGCAGCCATG	CGGAAATCAA	CCGTCTGATG	GCGATTTTGC	AAAAACAGGG	1260
CTGCCGCGGC	GTGGTCGGGA	TCGGCGGTGG	TAAAACCCTC	GATACCGCGA	AGGCGATCGG	1320
TTACTACCAG	AAGCTGCCGG	TGGTGGTGAT	CCCGACCATC	GCCTCGACCG	ATGCGCCAAC	1380

WO 96/3	5796				PCT/	US96/06705
CAGCGCGCTG	TCGGTGATCT	ACACCGAAGC	GGGCGAGTTT	GAAGAGTATC	TGATCTATCC	1440
GAAAAACCCG	GATATGGTGG	TGATGGACAC	GGCGATTATC	GCCAAAGCGC	CGGTACGCCT	1500
GCTGGTCTCC	GGCATGGGCG	ATGCGCTCTC	CACCTGGTTC	GAGGCCAAAG	CTTGCTACGA	1560
TGCGCGCGCC	ACCAGCATGG	CCGGAGGACA	GTCCACCGAG	GCGGCGCTGA	GCCTCGCCCG	1620
CCTGTGCTAT	GATACGCTGC	TGGCGGAGGG	CGAAAAGGCC	CGTCTGGCGG	CGCAGGCCGG	1680
GGTAGTGACC	GAAGCGCTGG	AGCGCATCAT	CGAGGCGAAC	ACTTACCTCA	GCGGCATTGG	1740
CTTTGAAAGC	AGTGGCCTGG	CCGCTGCCCA	TGCAATCCAC	AACGGTTTCA	CCATTCTTGA	1800
AGAGTGCCAT	CACCTGTATC	ACGGTGAGAA	AGTGGCCTTC	GGTACCCTGG	CGCAGCTGGT	1860
GCTGCAGAAC	AGCCCGATGG	ACGAGATTGA	AACGGTGCAG	GGCTTCTGCC	AGCGCGTCGG	1920
CCTGCCGGTG	ACGCTCGCGC	AGATGGGCGT	CAAAGAGGGG	ATCGACGAGA	AAATCGCCGC	1980
GGTGGCGAAA	GCTACCTGCG	CGGAAGGGGA	AACCATCCAT	AATATGCCGT	TTGCGGTGAC	2040
CCCGGAGAGC	GTCCATGCCG	CTATCCTCAC	CGCCGATCTG	TTAGGCCAGC	AGTGGCTGGC	2100
GCGTTAATTC	GCGGTGGCTA	AACCGCTGGC	CCAGGTCAGC	GGTTTTTCTT	TCTCCCCTCC	2160
GGCAGTCGCT	GCCGGAGGGG	TTCTCTATGG	TACAACGCGG	AAAAGGATAT	GACTGTTCAG	2220
ACTCAGGATA	CCGGGAAGGC	GGTCTCTTCC	GTCATTGCCC	AGTCATGGCA	CCGCTGCAGC	2280
AAGTTTATGC	AGCGCGAAAC	CTGGCAAACG	CCGCACCAGG	CCCAGGGCCT	GACCTTCGAC	2340
TCCATCTGTC	GGCGTAAAAC	CGCGCTGCTC	ACCATCGGCC	AGGCGGCGCT	GGAAGACGCC	2400
TGGGAGTTTA	TGGACGGCCG	CCCCTGCGCG	CTGTTTATTC	TTGATGAGTC	CGCCTGCATC	2460
CTGAGCCGTT	GCGGCGAGCC	GCAAACCCTG	GCCCAGCTGG	CTGCCCTGGG	ATTTCGCGAC	2520
GGCAGCTATT	GTGCGGAGAG	CATTATCGGC	ACCTGCGCGC	TGTCGCTGGC	CGCGATGCAG	2580
GGCCAGCCGA	TCAACACCGC	CGGCGATCGG	CATTTTAAGC	AGGCGCTACA	GCCATGGAGT	2640
TTTTGCTCGA	CGCCGGTGTT	TGATAACCAC	GGGCGGCTGT	TCGGCTCTAT	CTCGCTTTGC	2700
TGTCTGGTCG	AGCACCAGTC	CAGCGCCGAC	CTCTCCCTGA	CGCTGGCCAT	CGCCCGCGAG	2760
GTGGGTAACT	CCCTGCTTAC	CGACAGCCTG	CTGGCGGAAT	CCAACCGTCA	CCTCAATCAG	2820
ATGTACGGCC	TGCTGGAGAG	CATGGACGAT	GGGGTGATGG	CGTGGAACGA	ACAGGGCGTG	2880
CTGCAGTTTC	TCAATGTTCA	GGCGGCGAGA	CTGCTGCATC	TTGATGCTCA	GGCCAGCCAG	2940
GGGAAAAATA	TCGCCGATCT	GGTGACCCTC	CCGGCGCTGC	TGCGCCGCGC	CATCAAACAC	3000
GCCGCGGCC	TGAATCACGT	CGAAGTCACC	TTTGAAAGTC	AGCATCAGTT	TGTCGATGCG	3060
GTGATCACCT	TAAAACCGAT	TGTCGAGGCG	CAAGGCAACA	GTTTTATTCT	GCTGCTGCAT	3120

WO 96/35796

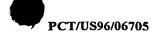
CCGGTGGAGC AGATGCGGCA GCTGATGACC AGCCAGCTCG GTAAAGTCAG CCACACCTTT 3180 GAGCAGATGT CTGCCGACGA TCCGGAAACC CGACGCCTGA TCCACTTTGG CCGCCAGGCG 3240 GCGCGCGCG GCTTCCCGGT GCTACTGTGC GGCGAAGAGG GGGTCGGGAA AGAGCTGCTG 3300 AGCCAGGCTA TTCACAATGA, AAGCGAACGG GCGGGCGGCC CCTACATCTC CGTCAACTGC 3360 CAGCTATATG CCGACAGCGT GCTGGGCCAG GACTTTATGG GCAGCGCCCC TACCGACGAT 3420 GAAAATGGTC GCCTGAGCCG CCTTGAGCTG GCCAACGGCG GCACCCTGTT TCTGGAAAAG 3480 ATCGAGTATC TGGCGCCGGA GCTGCAGTCG GCTCTGCTGC AGGTGATTAA GCAGGGCGTG 3540 CTCACCCGCC TCGACGCCCG GCGCCTGATC CCGGTGGATG TGAAGGTGAT TGCCACCACC 3600 ACCGTCGATC TGGCCAATCT GGTGGAACAG AACCGCTTTA GCCGCCAGCT GTACTATGCG 3660 CTGCACTCCT TTGAGATCGT CATCCCGCCG CTGCGCGCCC GACGCAACAG TATTCCGTCG 3720 CTGGTGCATA ACCGGTTGAA GAGCCTGGAG AAGCGTTTCT CTTCGCGACT GAAAGTGGAC 3780 GATGACGCGC TGGCACAGCT GGTGGCCTAC TCGTGGCCGG GGAATGATTT TGAGCTCAAC 3840 AGCGTCATTG AGAATATCGC CATCAGCAGC GACAACGGCC ACATTCGCCT GAGTAATCTG 3900 CCGGAATATC TCTTTTCCGA GCGGCCGGGC GGGGATAGCG CGTCATCGCT GCTGCCGGCC 3960 AGCCTGACTT TTAGCGCCAT CGAAAAGGAA GCTATTATTC ACGCCGCCCG GGTGACCAGC 4020 GGGCGGGTGC AGGAGATGTC GCAGCTGCTC AATATCGGCC GCACCACCCT GTGGCGCAAA 4080 ATGAAGCAGT ACGATATTGA CGCCAGCCAG TTCAAGCGCA AGCATCAGGC CTAGTCTCTT 4140 CGATTCGCGC CATGGAGAAC AGGGCATCCG ACAGGCGATT GCTGTAGCGT TTGAGCGCGT 4200 CGCGCAGCGG ATGCGCGCGG TCCATGGCCG TCAGCAGGCG TTCGAGCCGA CGGGACTGGG 4260 TGCGCGCCAC GTGCAGCTGG GCAGAGGCGA GATTCCTCCC CGGGATCACG AACTGTTTTA 4320 ACGGGCCGCT CTCGGCCATA TTGCGGTCGA TAAGCCGCTC CAGGGCGGTG ATCTCCTCTT 4380 CGCCGATCGT CTGGCTCAGG CGGGTCAGGC CCCGCGCATC GCTGGCCAGT TCAGCCCCCA 4440 GCACGAACAG CGTCTGCTGA ATATGGTGCA GGCTTTCCCG CAGCCCGGCG TCGCGGGTCG 4500 TGGCGTAGCA GACGCCCAGC TGGGATATCA GTTCATCGAC GGTGCCGTAG GCCTCGACGC 4560 GAATATGGTC TTTCTCGATG CGGCTGCCGC CGTACAGGGC GGTGGTGCCT TTATCCCCGG 4620 TGCGGGTATA GATACGATAC ATTCAGTTTC TCTCACTTAA CGGCAGGACT TTAACCAGCT 4680 GCCCGGCGTT GGCGCCGAGC GTACGCAGTT GATCGTCGCT ATCGGTGACG TGTCCGGTAG 4740 4800 CCAGCGCGC GTCCGCCGC AGCTGGGCAT GAGTGAGGGC TATCTCGCCG GACGCGCTGA GCCCGATACC CACCCGCAGG GGCGAGCTTC TGGCCGCCAG GGCGCCCAGC GCAGCGGCGT 4860

PCT/US96/06705

	WO 96/35	5796				PC	r/US96/06705
	CACCGCCTCC	GTCATAGGTT	ATGGTCTGGC	AGGGGACCCC	CTGCTCCTCC	AGCCCCCAGC	4920
	ACAGCTCATT	GATGGCGCCG	GCATGGTGCC	CGCGCGGATC	GTAAAACAGG	CGTACGCCTG	4980
	GCGGTGAAAG	CGACATGACG	GTCCCCTCGT	TAACACTCAG	AATGCCTGGC	GGAAAATCGC	5040
	GGCAATCTCC	TGCTCGTTGC	CTTTACGCGG	GTTCGAGAAC	GCATTGCCGT	CTTTTAGAGC	5100
	CATCTCCGCC	ATGTAGGGGA	AGTCGGCCTC	TTTTACCCCC	AGATCGCGCA	GATGCTGCGG	5160
	AATACCGATA	TCCATCGACA	GACGCGTGAT	AGCGGCGATG	GCTTTTTCCG	CCGCGTCGAG	5220
	AGTGGACAGT	CCGGTGATAT	TTTCGCCCAT	CAGTTCAGCG	ATATCGGCGA	ATTTCTCCGG	5280
	GTTGGCGATC	AGGTTGTAGC	GCGCCACATG	CGGCAGCAGG	ACAGCGTTGG	CCACGCCGTG	5340
	CGGCATGTCG	TACAGGCCGC	CCAGCTGGTG	CGCCATGGCG	TGCACGTAGC	CGAGGTTGGC	5400
	GTTATTGAAA	GCCATCCCGG	CCAGCAGAGA	AGCATAGGCC	ATGTTTTCCC	GCGCCTGCAG	5460
	ATTGCTGCCG	AGGGCCACGG	CCTGGCGCAG	GTTGCGGGCG	ATGAGGCGGA	TCGCCTGCAT	5520
122	Y .	TCCGTCACCG	GGTTAGCGTC	TTTGGAGATA	TAGGCCTCTA	CGGCGTGGGT	5580
ŭ.	CAGGGCATCC	ATCCCGGTCG	CCGCGGTCAG	GGCGGCCGGT	TTACCGATCA	TCAGCAGTGG	5640
	ATCGTTGATA	GAGACCGACG	GCAGTTTGCG	CCAGCTGACG	ATCACAAACT	TCACTTTGGT	5700
	TTCGGTGTTG	GTCAGGACGC	AGTGGCGGGT	GACCTCGCTG	GCGGTGCCGG	CGGTGGTATT	5760
100	;	ATAGGCGGCA	GCGGGTTGGT	CAGGGTCTCG	ATTCCGGCAT	ACTGGTACAG	5820
		TGGGTGGCGG	CGATGCCGAT	GCCTTTGCCG	CAATCGTGCG	GGCTGCCGCC	5880
FL.	GCCCACGGTG	ACGATGATGT	CGCACTGTTC	GCGGCGAAAC	ACGGCGAGGC	CGTCGCGCAC	5940
	GTTGGTGTCT	TTCGGGTTCG	GCTCGACGCC	GTCAAAGATC	GCCACCTCGA	TCCCGGCCTC	6000
بير	CCGCAGATAA	TGCAGGGTTT	TGTCCACCGC	GCCATCTTTA	ATTGCCCGCA	GGCCTTTGTC	6060
	GGTGACCAGC	AGGGCTTTTT	TCCCCCCAG	CAGCTGGCAG	CGTTCGCCGA	CTACGGAAAT	6120
	GGCGTTGGGG	CCAAAAAAGT	TAACGTTTGG	CACCAGATAA	TCAAACATAC	GATAGCTCAT	6180
	AATATACCTT	CTCGCTTCAG	GTTATAATGC	GGAAAAACAA	TCCAGGGCGC	ACTGGGCTAA	6240
	TAATTGATCC	TGCTCGACCG	TACCGCCGCT	AACGCCGACG	GCGCCAATTA	CCTGCTCATT	6300
	AAAAATAACT	GGCAGGCCGC	CGCCAAAAAT	AATAATTCGC	TGTTGGTTGG	TTAGCTGCAG	6360
	ACCGTACAGA	GATTGTCCTG	GCTGGACCGC	TGACGTAATT	TCATGGGTAC	CTTGCTTCAG	6420
	GCTGCAGGCG	CTCCAGGCTT	TATTCAGGGA	AATATCGCAG	CTGGAGACGA	AGGCCTCGTC	6480
•	CATCCGCTGG	ATAAGCAGCG	TGTTGCCTCC	GCGGTCAACT	ACGGAAAACA	CCACCGCCAC	6540
_	GTTGATCTCA	GTGGCTTTTT	TTTCCACCGC	CGCCGCCATT	TGCTGGGCGG	CGGCCAGGGT	6600

WO 96/35	WO 96/35796 PCT/US96/06705					
GATTGTCTGA	ACTTGTTGGC	TCTTGTTCAT	CATTCTCTCC	CGCACCAGGA	TAACGCTGGC	6660
GCGAATAGTC	AGTAGGGGGC	GATAGTAAAA	AACTATTACC	ATTCGGTTGG	CTTGCTTTAT	6720
TTTTGTCAGC	GTTATTTTGT	CGCCCGCCAT	GATTTAGTCA	ATAGGGTTAA	AATAGCGTCG	6780
GAAAAACGTA	ATTAAGGGCG	TTTTTTATTA	ATTGATTTAT	ATCATTGCGG	GCGATCACAT	6840
TTTTTATTTT	TGCCGCCGGA	GTAAAGTTTC	ATAGTGAAAC	TGTCGGTAGA	TTTCGTGTGC	6900
CAAATTGAAA	CGAAATTAAA	TTTATTTTTT	TCACCACTGG	CTCATTTAAA	GTTCCGCTAT	6960
TGCCGGTAAT	GGCCGGGCGG	CAACGACGCT	GGCCCGGCGT	ATTCGCTACC	GTCTGCGGAT	7020
TTCACCTTTT	GAGCCGATGA	ACAATGAAAA	GATCAAAACG	ATTTGCAGTA	CTGGCCCAGC	7080
GCCCCGTCAA	TCAGGACGGG	CTGATTGGCG	AGTGGCCTGA	AGAGGGGCTG	ATCGCCATGG	7140
ACAGCCCCTT	TGACCCGGTC	TCTTCAGTAA	AAGTGGACAA	CGGTCTGATC	GTCGAACTGG	7200
ACGGCAAACG	CCGGGACCAG	TTTGACATGA	TCGACCGATT	TATCGCCGAT	TACGCGATCA	7260
ACGTTGAGCG	CACAGAGCAG	GCAATGCGCC	TGGAGGCGGT	GGAAATAGCC	CGTATGCTGG	7320
TGGATATTCA	CGTCAGCCGG	GAGGAGATCA	TTGCCATCAC	TACCGCCATC	ACGCCGGCCA	7380
AAGCGGTCGA	GGTGATGGCG	CAGATGAACG	TGGTGGAGAT	GATGATGGCG	CTGCAGAAGA	7440
TGCGTGCCCG	CCGGACCCCC	TCCAACCAGT	GCCACGTCAC	CAATCTCAAA	GATAATCCGG	7500
TGCAGATTGC	CGCTGACGCC	GCCGAGGCCG	GGATCCGCGG	CTTCTCAGAA	CAGGAGACCA	7560
CGGTCGGTAT	CGCGCGCTAC	GCGCCGTTTA	ACGCCCTGGC	GCTGTTGGTC	GGTTCGCAGT	7620
GCGGCCGCCC	CGGCGTGTTG	ACGCAGTGCT	CGGTGGAAGA	GGCCACCGAG	CTGGAGCTGG	7680
GCATGCGTGG	CTTAACCAGC	TACGCCGAGA	CGGTGTCGGT	CTACGGCACC	GAAGCGGTAT	7740
TTACCGACGG	CGATGATACG	CCGTGGTCAA	AGGCGTTCCT	CGCCTCGGCC	TACGCCTCCC	7800
GCGGGTTGAA	AATGCGCTAC	ACCTCCGGCA	CCGGATCCGA	AGCGCTGATG	GGCTATTCGG	7860
AGAGCAAGTC	GATGCTCTAC	CTCGAATCGC	GCTGCATCTT	CATTACTAAA	GCCCCGGGG	7920
TTCAGGGACT	GCAAAACGGC	GCGGTGAGCT	GTATCGGCAT	GACCGGCGCT	GTGCCGTCGG	7980
GCATTCGGGC	GGTGCTGGCG	GAAAACCTGA	TCGCCTCTAT	GCTCGACCTC	GAAGTGGCGT	8040
CCGCCAACGA	CCAGACTTTC	TCCCACTCGG	ATATTCGCCG	CACCGCGCGC	ACCCTGATGC	8100
AGATGCTGCC	GGGCACCGAC	TTTATTTTCT	CCGGCTACAG	CGCGGTGCCG	AACTACGACA	8160
ACATGTTCGC	CGGCTCGAAC	TTCGATGCGG	AAGATTTTGA	TGATTACAAC	ATCCTGCAGC	8220
GTGACCTGAT	GGTTGACGGC	GGCCTGCGTC	CGGTGACCGA	GGCGGAAACC	ATTGCCATTC	8280
GCCAGAAAGC	GGCGCGGCG	ATCCAGGCGG	TTTTCCGCGA	GCTGGGGCTG	CCGCCAATCG	8340

CCGACGAGGA GGTGGAGGCC GCCACCTACG CGCACGGCAG CAACGAGATG CCGCCGCGTA	8400
ACGTGGTGGA GGATCTGAGT GCGGTGGAAG AGATGATGAA GCGCAACATC ACCGGCCTCG	8460
ATATTGTCGG CGCGCTGAGC CGCAGCGGCT TTGAGGATAT CGCCAGCAAT ATTCTCAATA	8520
TGCTGCGCCA GCGGGTCACC GGCGATTACC TGCAGACCTC GGCCATTCTC GATCGGCAGT	8580
TCGAGGTGGT GAGTGCGGTC AACGACATCA ATGACTATCA GGGGCCGGGC ACCGGCTATC	8640
GCATCTCTGC CGAACGCTGG GCGGAGATCA AAAATATTCC GGGCGTGGTT CAGCCCGACA	8700
CCATTGAATA AGGCGGTATT CCTGTGCAAC AGACAACCCA AATTCAGCCC TCTTTTACCC	8760
TGAAAACCCG CGAGGGCGGG GTAGCTTCTG CCGATGAACG CGCCGATGAA GTGGTGATCG	8820
GCGTCGGCCC TGCCTTCGAT AAACACCAGC ATCACACTCT GATCGATATG CCCCATGGCG	8880
CGATCCTCAA AGAGCTGATT GCCGGGGTGG AAGAAGAGGG GCTTCACGCC CGGGTGGTGC	8940
GCATTCTGCG CACGTCCGAC GTCTCCTTTA TGGCCTGGGA TGCGGCCAAC CTGAGCGGCT	9000
CGGGGATCGG CATCGGTATC CAGTCGAAGG GGACCACGGT CATCCATCAG CGCGATCTGC	9060
TGCCGCTCAG CAACCTGGAG CTGTTCTCCC AGGCGCCGCT GCTGACGCTG GAGACCTACC	9120
GGCAGATTGG CAAAAACGCT GCGCGCTATG CGCGCAAAGA GTCACCTTCG CCGGTGCCGG	9180
TGGTGAACGA TCAGATGGTG CGGCCGAAAT TTATGGCCAA AGCCGCGCTA TTTCATACA	9240
AAGAGACCAA ACATGTGGTG CAGGACGCCG AGCCCGTCAC CCTGCACATC GACTTAGTAA	9300
GGGAGTGACC ATGAGCGAGA AAACCATGCG CGTGCAGGAT TATCCGTTAG CCACCCGCTG	9360
CCCGGAGCAT ATCCTGACGC CTACCGGCAA ACCATTGACC GATATTACCC TCGAGAAGGT	9420
GCTCTCTGGC GAGGTGGGCC CGCAGGATGT GCGGATCTCC CGCCAGACCC TTGAGTACCA	9480
GGCGCAGATT GCCGAGCAGA TGCAGCGCCA TGCGGTGGCG CGCAATTTCC GCCGCGCGCGC	9540
GGAGCTTATC GCCATTCCTG ACGAGCGCAT TCTGGCTATC TATAACGCGC TGCGCCCGTT	9600
CCGCTCCTCG CAGGCGGAGC TGCTGGCGAT CGCCGACGAG CTGGAGCACA CCTGGCATGC	9660
GACAGTGAAT GCCGCCTTTG TCCGGGAGTC GGCGGAAGTG TATCAGCAGC GGCATAAGCT	9720
GCGTAAAGGA AGCTAAGCGG AGGTCAGCAT GCCGTTAATA GCCGGGATTG ATATCGGCAA	9780
CGCCACCACC GAGGTGGCGC TGGCGTCCGA CTACCCGCAG GCGAGGGCGT TTGTTGCCAG	9840
CGGGATCGTC GCGACGACGG GCATGAAAGG GACGCGGGAC AATATCGCCG GGACCCTCGC	9900
CGCGCTGGAG CAGGCCCTGG CGAAAACACC GTGGTCGATG AGCGATGTCT CTCGCATCTA	9960
TCTTAACGAA GCCGCGCGG TGATTGGCGA TGTGGCGATG GAGACCATCA CCGAGACCAT 1	.0020
TATCACCGAA TCGACCATGA TCGGTCATAA CCCGCAGACG CCGGGCGGGG TGGGCGTTGG 1	0800



CGTGGGGACG	ACTATCGCCC	TCGGGCGGCT	GGCGACGCTG	CCGGCGGCGC	AGTATGCCGA	10140
GGGGTGGATC	GTACTGATTG	ACGACGCCGT	CGATTTCCTT	GACGCCGTGT	GGTGGCTCAA	10200
TGAGGCGCTC	GACCGGGGGA	TCAACGTGGT	GGCGGCGATC	CTCAAAAAGG	ACGACGGCGT	10260
GCTGGTGAAC	AACCGCCTGC	GTAAAACCCT	GCCGGTGGTG	GATGAAGTGA	CGCTGCTGGA	10320
GCAGGTCCCC	GAGGGGGTAA	TGGCGGCGGT	GGAAGTGGCC	ececceecc	AGGTGGTGCG	10380
GATCCTGTCG	AATCCCTACG	GGATCGCCAC	CTTCTTCGGG	CTAAGCCCGG	AAGAGACCCA	10440
GGCCATCGTC	CCCATCGCCC	GCGCCCTGAT	TGGCAACCGT	TCCGCGGTGG	TGCTCAAGAC	10500
CCCGCAGGGG	GATGTGCAGT	CGCGGGTGAT	CCCGGCGGGC	AACCTCTACA	TTAGCGGCGA	10560
AAAGCGCCGC	GGAGAGGCCG	ATGTCGCCGA	GGGCGCGGAA	GCCATCATGC	AGGCGATGAG	10620
CGCCTGCGCT	CCGGTACGCG	ACATCCGCGG	CGAACCGGGC	ACCCACGCCG	GCGGCATGCT	10680
TGAGCGGGTG	CGCAAGGTAA	TGGCGTCCCT	GACCGGCCAT	GAGATGAGCG	CGATATACAT	10740
CCAGGATCTG	CTGGCGGTGG	ATACGTTTAT	TCCGCGCAAG	GTGCAGGGCG	GGATGGCCGG	10800
CGAGTGCGCC	ATGGAGAATG	CCGTCGGGAT	GGCGGCGATG	GTGAAAGCGG	ATCGTCTGCA	10860
AATGCAGGTT	ATCGCCCGCG	AACTGAGCGC	CCGACTGCAG	ACCGAGGTGG	TGGTGGGCGG	10920
CGTGGAGGCC	AACATGGCCA	TCGCCGGGGC	GTTAACCACT	CCCGGCTGTG	CGGCGCCGCT	10980
GGCGATCCTC	GACCTCGGCG	CCGGCTCGAC	GGATGCGGCG	ATCGTCAACG	CGGAGGGGCA	11040
GATAACGGCG	GTCCATCTCG	cceeecee	GAATATGGTC	AGCCTGTTGA	TTAAAACCGA	11100
GCTGGGCCTC	GAGGATCTTT	CGCTGGCGGA	AGCGATAAAA	AAATACCCGC	TGGCCAAAGT	11160
GGAAAGCCTG	TTCAGTATTC	GTCACGAGAA	TGGCGCGGTG	GAGTTCTTTC	GGGAAGCCCT	11220
CAGCCCGGCG	GTGTTCGCCA	AAGTGGTGTA	CATCAAGGAG	GGCGAACTGG	TGCCGATCGA	11280
TAACGCCAGC	CCGCTGGAAA	AAATTCGTCT	CGTGCGCCGG	CAGGCGAAAG	AGAAAGTGTT	11340
TGTCACCAAC	TGCCTGCGCG	CGCTGCGCCA	GGTCTCACCC	GGCGGTTCCA	TTCGCGATAT	11400
CGCCTTTGTG	GTGCTGGTGG	GCGGCTCATC	GCTGGACTTT	GAGATCCCGC	AGCTTATCAC	11460
GGAAGCCTTG	TCGCACTATG	GCGTGGTCGC	CGGGCAGGGC	AATATTCGGG	GAACAGAAGG	11520
GCCGCGCAAT	GCGGTCGCCA	CCGGGCTGCT	ACTGGCCGGT	CAGGCGAATT	AAACGGGCGC	11580
TCGCGCCAGC	CTCTCTCTTT	AACGTGCTAT	TTCAGGATGC	CGATAATGAA	CCAGACTTCT	11640
ACCTTAACCG	GGCAGTGCGT	GGCCGAGTTT	CTTGGCACCG	GATTGCTCAT	TTTCTTCGGC	11700
GCGGGCTGCG	TCGCTGCGCT	GCGGGTCGCC	GGGGCCAGCT	TTGGTCAGTG	GGAGATCAGT	11760
ATTATCTGGG	GCCTTGGCGT	CGCCATGGCC	ATCTACCTGA	CGGCCGGTGT	CTCCGGCGCG	11820

WO 96/35	796 PC	r/US96/06705
CACCTAAATC	CGGCGGTGAC CATTGCCCTG TGGCTGTTCG CCTGTTTTGA ACGCCGCAAG	11880
GTGCTGCCGT	TTATTGTTGC CCAGACGGCC GGGGCCTTCT GCGCCGCCGC GCTGGTGTAT	11940
GGGCTCTATC	GCCAGCTGTT TCTCGATCTT GAACAGAGTC AGCATATCGT GCGCGGCACT	12000
GCCGCCAGTC	TTAACCTGGC CGGGGTCTTT TCCACGTACC CGCATCCACA TATCACTTTT	12060
ATACAAGCGT	TTGCCGTGGA GACCACCATC ACGGCAATCC TGATGGCGAT GATCATGGCC	12120
CTGACCGACG	ACGGCAACGG AATTC	12145
(2)	INFORMATION FOR SEQ ID NO:2:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
	ATTCAT GAAAAGATCA AAACGATTTG	30
(2)	~ · ·	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GCGA	ATTCTT ATTCAATGGT GTCGGGCTG	29
(2)	INFORMATION FOR SEQ ID NO:4:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GCGA	ATTCAT GCAACAGACA ACCCAAATTC	30
(2)	INFORMATION FOR SEQ ID NO: 5:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 25 base pairs	

WO 96/35796		PCT/US96/06705
	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:	
GCGAATTCAC	TCCCTTACTA AGTCG	25
(2) INFOR	RMATION FOR SEQ ID NO: 6:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	
GCGAATTCAT	GAGCTATCGT ATGTTTG	27
(2) INFOR	RMATION FOR SEQ ID NO:7:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
GCGAATTCAG	AATGCCTGGC GGAAAATC	28
(2) INFOR	RMATION FOR SEQ ID NO:8:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA (genomic)	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:8:	
GGGAATTCAT	GAGCGAGAAA ACCATGCG	28
(2) INFOR	RMATION FOR SEQ ID NO:9:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid	

WO 96/35796

(C) STRANDEDNESS: single (D) TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) (ii) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: GCGAATTCTT AGCTTCCTTT ACGCAGC 27 (2) INFORMATION FOR SEQ ID NO:10: SEQUENCE CHARACTERISTICS: (A) LENGTH: 94 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO:10: AGCTTAGGAG TCTAGAATAT TGAGCTCGAA TTCCCGGGCA TGCGGTACCG GATCCAGAAA 60 AAAGCCCGCA CCTGACAGTG CGGGCTTTTT TTTT 94 INFORMATION FOR SEQ ID NO:11: (2) SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 33 GGCCAAGCTT AAGGAGGTTA ATTAAATGAA AAG (2) INFORMATION FOR SEQ ID NO:12: SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO:12: (xi) 26 GCTCTAGATT ATTCAATGGT GTCGGG INFORMATION FOR SEQ ID NO:13: (2) SEQUENCE CHARACTERISTICS: (A) LENGTH: 42 base pairs (B) TYPE: nucleic acid

PCT/US96/06705

WO 96/35796 PCT/US96/06705 (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii)MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO:13: (xi) GCGCCGTCTA GAATTATGAG CTATCGTATG TTTGATTATC TG 42 (2) INFORMATION FOR SEQ ID NO:14: SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO:14: TCTGATACGG GATCCTCAGA ATGCCTGGCG GAAAAT 36 (2) INFORMATION FOR SEQ ID NO:15: SEQUENCE CHARACTERISTICS: (A) LENGTH: 181 base pairs (B) TYPE: nucleic acid STRANDEDNESS: single (C) (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO:15: CGATCTGTGC TGTTTGCCAC GGTATGCAGC ACCAGCGCGA GATTATGGGC TCGCACGCTC 60 GACTGTCGGA CGGGGGCACT GGAACGAGAA GTCAGGCGAG CCGTCACGCC CTTGACAATG 120 CCACATCCTG AGCAAATAAT TCAACCACTA AACAAATCAA CCGCGTTTCC CGGAGGTAAC 180 С 181 (2) INFORMATION FOR SEQ ID NO:16: SEQUENCE CHARACTERISTICS: (A) LENGTH: 149 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16: CGATCTGTGC TGTTTGCCAC GGTATGCAGC ACCAGCGCGA GATTATGGGC TCGCACGCTC 60

120

GACTGTCGGA CGGGGGCACT GGAACATGCC ACATCCTGAG CAAATAATTC AACCACTAAA

WO 96/35796 PCT/US96/06705 CAAATCAACC GCGTTTCCCG GAGGTAACC 149 INFORMATION FOR SEQ ID NO:17: SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17: GGAATTCACT AGTCGATCTG TGCTGTTTGC CAC 33 INFORMATION FOR SEQ ID NO:18: (2) SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18: GGGGAAGCTT GGTTACCTCC GGGAAACGCG GTT 33 INFORMATION FOR SEQ ID NO:19: (2) SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19: TCGACCACAA GGAGGA 16 INFORMATION FOR SEQ ID NO:20: (2) SEQUENCE CHARACTERISTICS: (i)

(A) LENGTH: 16 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

CTAGTCCTCC TTGTGG

WO 96/35796 PCT/US96/06705 INFORMATION FOR SEQ ID NO:21: (2) SEQUENCE CHARACTERISTICS: (A) LENGTH: 45 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO:21: ACTGGCCGTC GTTTTACTCG AGTCGTGACT GGGAAAACCC TGGCG 45 INFORMATION FOR SEQ ID NO:22: (2) SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) SEQUENCE DESCRIPTION: SEQ ID NO:22: (xi) AATTCAAAGG AGGT 14 INFORMATION FOR SEQ ID NO:23: (2) SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 14 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) (ii) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23: CTAGACCTCC TTTG 14 INFORMATION FOR SEQ ID NO:24: (2) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) (ii) SEQUENCE DESCRIPTION: SEQ ID NO:24: (xi)

AGCTTGTCGA CCATGAAAA

19

WO 96/357	796	PCT/US96/06705
(2)	INFORMATION FOR SEQ ID NO:25:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:	
GATCT	TTTTCA TGGTCGACA	19
(2)	INFORMATION FOR SEQ ID NO:26:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 13 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:	
TCGAC	CCAGGA GGA	13
(2)	INFORMATION FOR SEQ ID NO:27:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 13 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:	
CTAGI	TCCTCC TGG	13
(2)	INFORMATION FOR SEQ ID NO:28:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:	
TCGAC	CGAATT CAGGAGGA	18

WO 96/35	5796	PCT/US96/06705
(2)	INFORMATION FOR SEQ ID NO:29:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29	:
CTAG	STECTEC TGAATTEG	18
(2)	INFORMATION FOR SEQ ID NO:30:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30	:
ATGT	ACAAGA TCCTGATCGC CGA	23
(2)	INFORMATION FOR SEQ ID NO:31:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31	:
TCAG	CGGCGC AGGTAGGCGG CG	22
(2)	INFORMATION FOR SEQ ID NO:32:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32	:
ልጥርል	CCAACG GCCGGATCCG TCGACCTGCA G	31

O 96/35796	PCT/US96/06705
(2) INFORMATION FOR SEQ ID NO:33:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:	
CTACCCTTGG CCCCGGATCC GTCGACCTGC AG	32
(2) INFORMATION FOR SEQ ID NO:34:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
CACGGCCTGG CGCAGGTTGC GGG	23
(2) INFORMATION FOR SEQ ID NO:35:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:	
GGCAGCCCGC ACGATTGCGG C	21
(2) INFORMATION FOR SEQ ID NO:36:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:	
GCGGAAAACC GCCTGGATCG C	21



(2) IN	FORMATION FOR SEQ ID NO:37:	
. ((i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
(i	i) MOLECULE TYPE: DNA (genomic)	
(x	i) SEQUENCE DESCRIPTION: SEQ ID NO:37:	
GGGTTCAG	GG ACTGCAAAAC G	21
(2) INE	FORMATION FOR SEQ ID NO:38:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 35 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i	i) MOLECULE TYPE: DNA (genomic)	
(x	i) SEQUENCE DESCRIPTION: SEQ ID NO:38:	
GGAATTCAG	GA TCTCAGCAAT GAAAAGATCA AAACG	35
(2) INF	FORMATION FOR SEQ ID NO:39:	
(I) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i	i) MOLECULE TYPE: DNA (genomic)	
(x	i) SEQUENCE DESCRIPTION: SEQ ID NO:39:	
GGAATTCAG	GA TCTCAGCAAT GCAACAGACA ACCC	34
(2) INF	FORMATION FOR SEQ ID NO:40:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(i	i) MOLECULE TYPE: DNA (genomic)	
(x	i) SEQUENCE DESCRIPTION: SEQ ID NO:40:	
GCTCTAGAT	TC ACTCCCCTTA CTAAGTCG	28

O 96/35796	PCT/US96/06705
(2) INFORMATION FOR SEQ ID NO:41:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 37 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(b) 1010B031. 11Meal	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:	
GGAATTCAGA TCTCAGCAAT GAGCGAGAAA ACCATGC	37
(2) INFORMATION FOR SEQ ID NO:42:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 27 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 42:	
GCTCTAGATT AGCTTCCTTT ACGCAGC	27
(2) INFORMATION FOR SEQ ID NO:43:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 38 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:	
GGAATTCAGA TCTCAGCAAT GAGCTATCGT ATGTTTGA	38
(2) INFORMATION FOR SEQ ID NO:44:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 24 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA (genomic)	
(II) NOBBOODD IIID. Dim (genomic)	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:	
GCTCTAGATC AGAATGCCTG GCGG	24



WO 96/35796



- (2) INFORMATION FOR SEQ ID NO:45:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 35 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:

GGAATTCAGA TCTAGCAATG CCGTTAATAG CCGGG

35

- (2) INFORMATION FOR SEQ ID NO:46:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:

GCTCTAGATT AATTCGCCTG ACCGGC

26